



App No.: 10/071,894

DOI: 10.1002/1522-2675(200209)24:09<1522-2675::AID-MOLB1522>3.0.CO;2-1

Title: Glyoxylate Cycle Enzymes As Targets...

Inventors: Michael C. Lorenz, *et al.*

Figure 1A

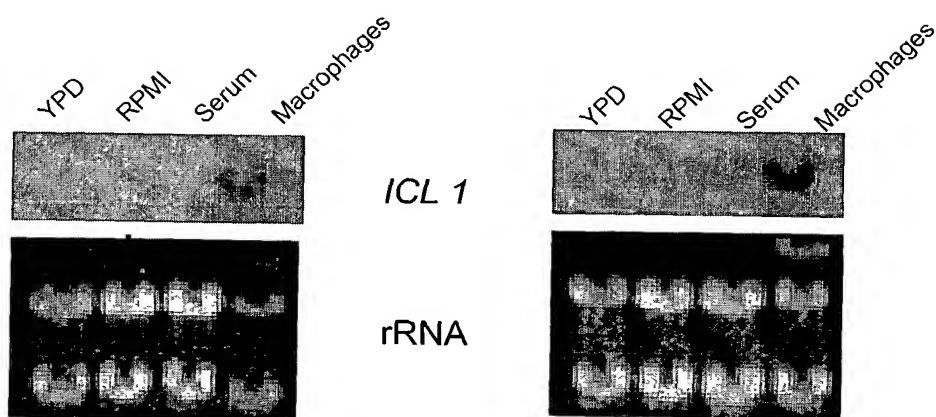
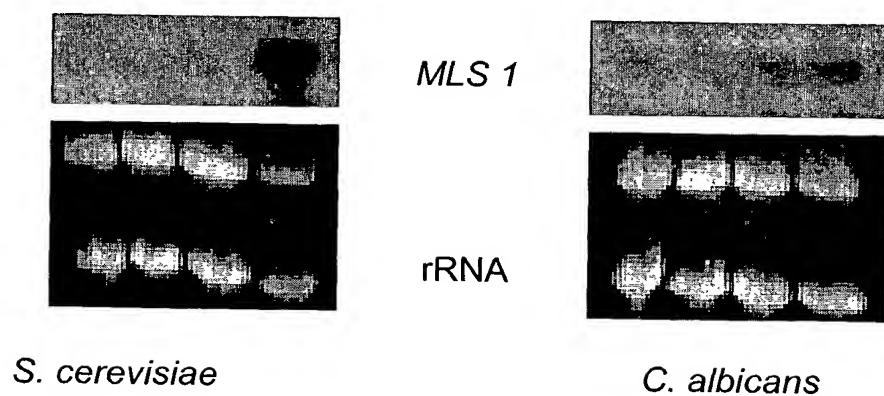
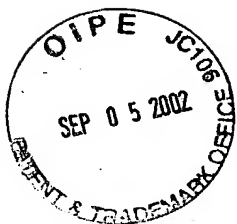


Figure 1B





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Figure 2

A

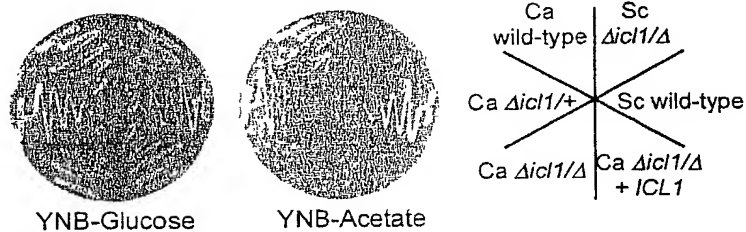
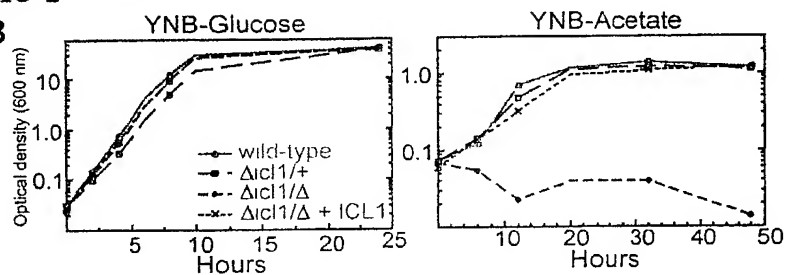


Figure 2

B





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Figure 3

A

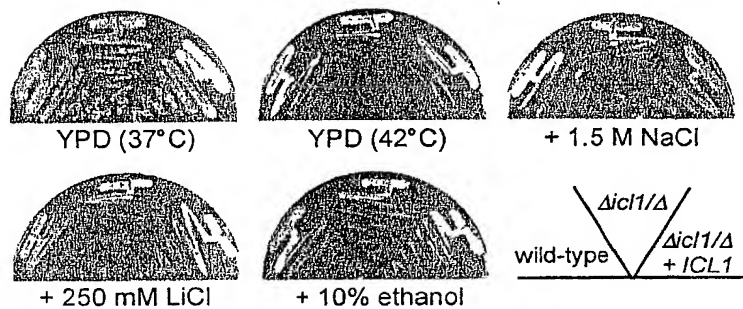


Figure 3

B

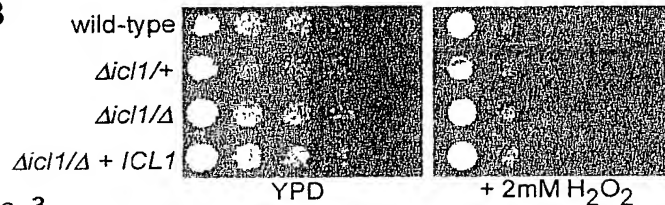
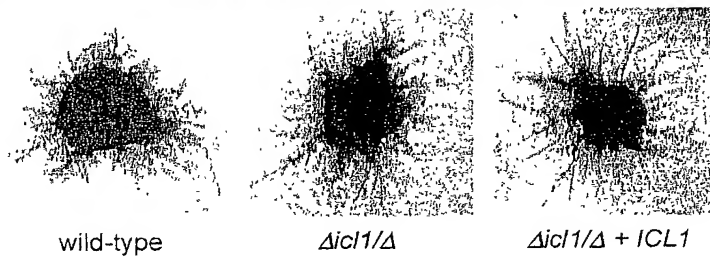


Figure 3

C





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Isocitrate lyase mutations attenuate virulence in *C. albicans*

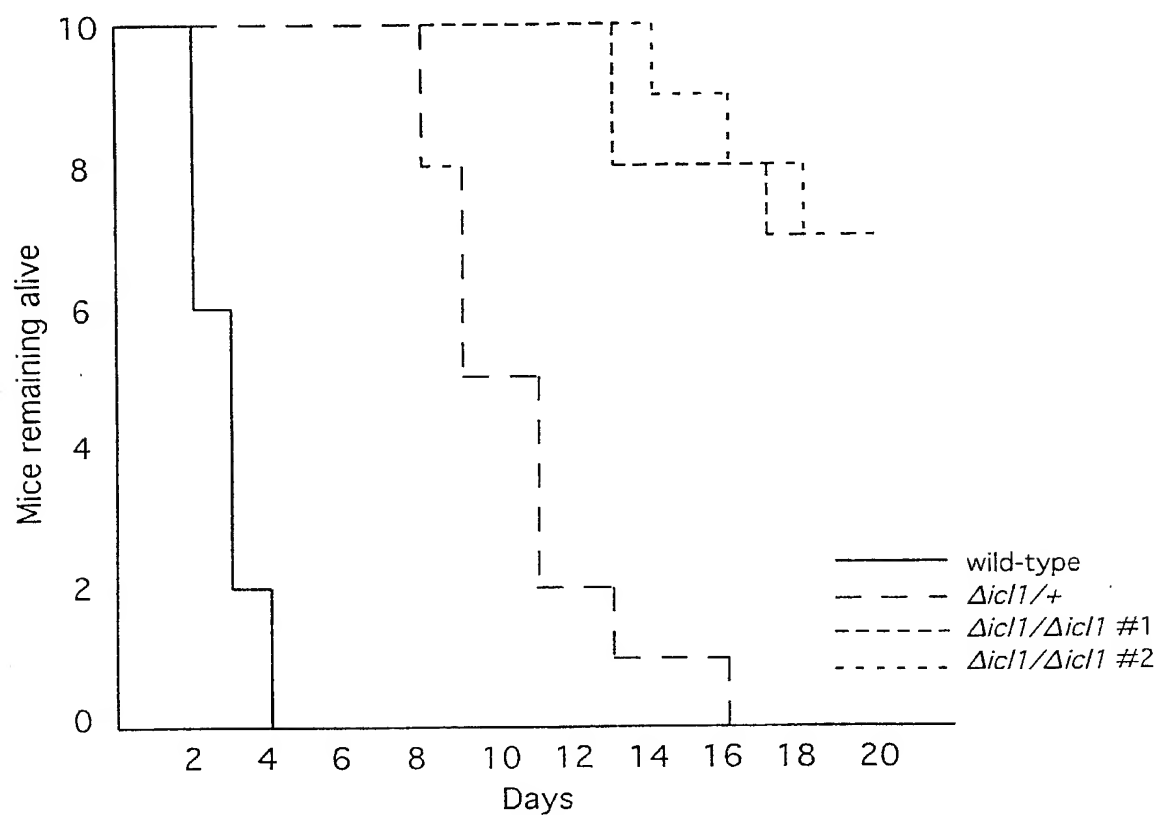


Figure 4

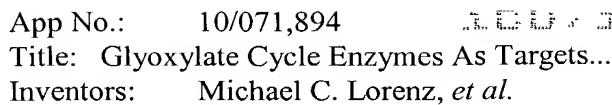


Figure 1 displays a 10x10 grid of 100 amino acid sequence logos. Each logo represents a position in a protein sequence, with the y-axis showing the frequency of each amino acid (A, C, G, U) and the x-axis showing the position (1-100). The logos are arranged in a grid, with each logo corresponding to a specific amino acid sequence. The logos are color-coded: A (blue), C (green), G (red), and U (yellow). The logos are arranged in a grid, with each logo corresponding to a specific amino acid sequence. The logos are color-coded: A (blue), C (green), G (red), and U (yellow).

Figure 5



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The regulation of *ICL1* is similar in both *S. cerevisiae* and *C. albicans*

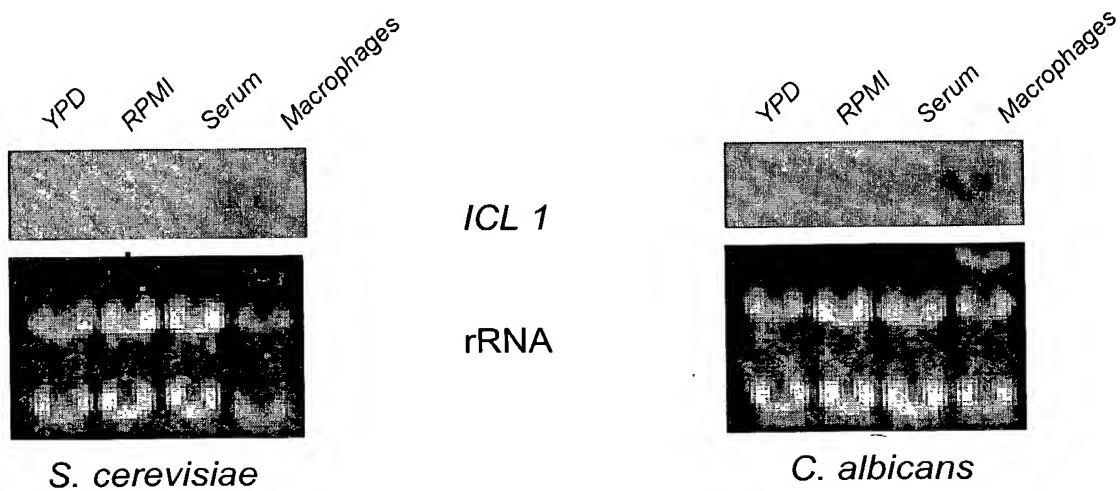


Figure 6

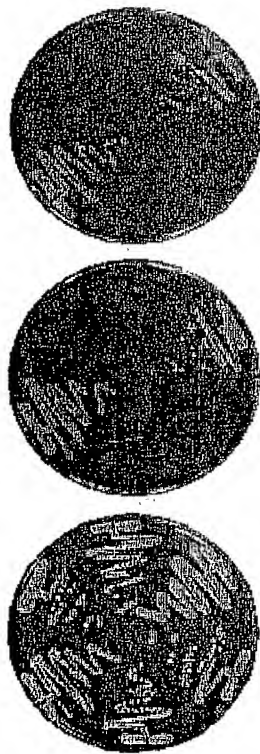
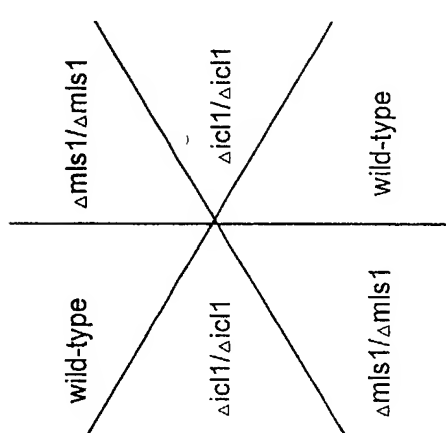


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YNB-2% Glucose YNB-2% Acetate YNB-2% Ethanol S. cerevisiae C. albicans

FIGURE 7

C. albicans glyoxylate mutants: Growth rates

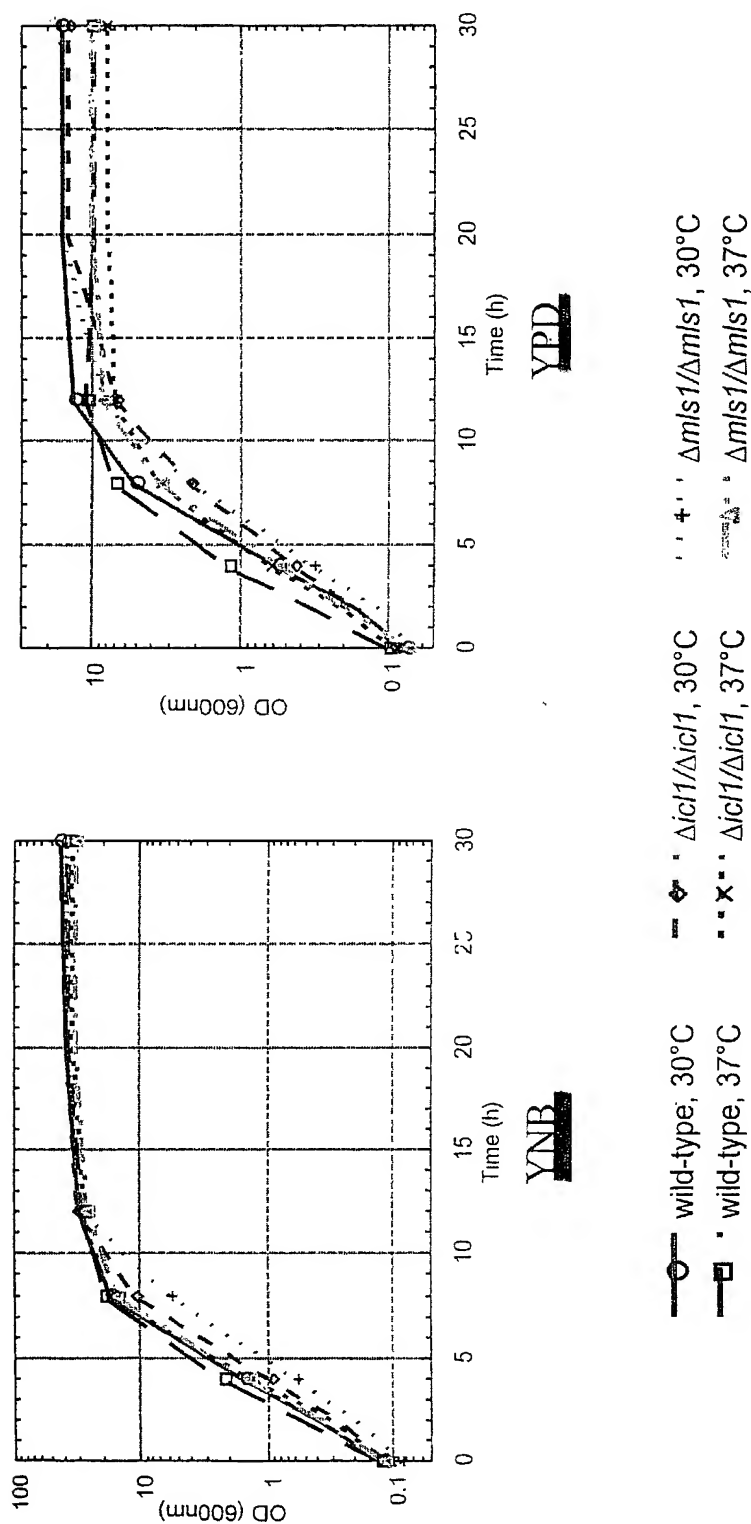


Figure 8

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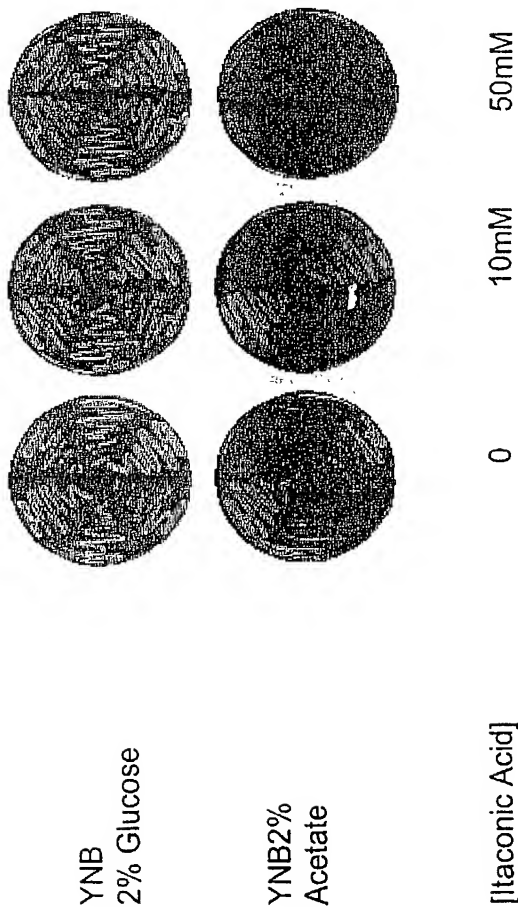
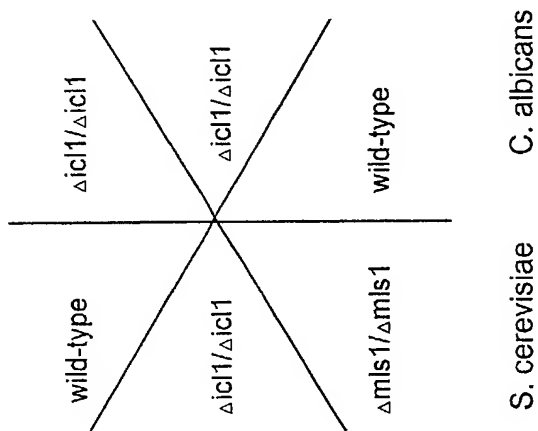


FIGURE 9